SEQUENCE LISTING

1

<110> Crooke, Stanley T Lima, Walter Wu, Hongjiang

<120> Methods of Using Mammalian RNase H and Compositions Thereof

<130> ISPH-0520

<140> US 09/781,712

<141> 2001-02-12

<150> US 60/067,458

<151> 1997-12-04

<150> US 09/203,716

<151> 1998-12-02

<150> US 09/343,809

<151> 1999-06-30

<150> US 09/684,254

<151> 2000-10-06

<160> 39

<170> PatentIn version 3.1

<210> 1

<211> 299

<212> PRT

<213> Homo sapien

<400> 1

Met Asp Leu Ser Glu Leu Glu Arg Asp Asn Thr Gly Arg Cys Arg Leu 1 5 10 15

Ser Ser Pro Val Pro Ala Val Cys Arg Lys Glu Pro Cys Val Leu Gly 20 25 30

Val Asp Glu Ala Gly Arg Gly Pro Val Leu Gly Pro Met Val Tyr Ala 35 40 45

Ile Cys Tyr Cys Pro Leu Pro Arg Leu Ala Asp Leu Glu Ala Leu Leu 50 55 60

Val Ala Asp Ser Leu Thr Leu Leu Glu Ser Glu Arg Glu Arg Leu Phe 65 70 75 80

Ala Leu Met Glu Asp Thr Asp Phe Val Gly Trp Ala Leu Asp Val Leu 85 90 95

Ser Pro Asn Leu Ile Ser Thr Ser Met Leu Gly Trp Val Leu Tyr Asn

100 105 110

Leu Asn Ser Leu Ser His Asp Thr Ala Thr Gly Leu Ile Gln Tyr Ala 115 120 125

Leu Asp Gln Gly Val Asn Val Thr Gln Val Phe Val Asp Thr Val Gly 130 135 140

Glu Val Thr Val Leu Ala Leu Ala Asp Ala Leu Tyr Pro Val Val Ser 165 170 175

Ala Ala Ser Ile Cys Ala Leu Val Ala Arg Asp Gln Ala Val Leu Leu 180 185 190

Trp Gln Phe Val Glu Leu Leu Gln Asp Leu Asp Thr Asp Tyr Gly Ser 195 200 205

Gly Tyr Pro Asn Asp Pro Leu Thr Leu Ala Trp Leu Leu Glu His Val 210 215 220

Glu Pro Val Phe Gly Phe Pro Gln Phe Val Arg Phe Ser Trp Arg Thr 225 230 235 240

Ala Gl
n Thr Ile Leu Glu Leu Glu Ala Glu Asp Val Ile Tr
p Glu Asp 245 250 255

Ser Ala Ser Glu Asn Gln Glu Gly Leu Arg Leu Ile Thr Ser Tyr Phe $260 \hspace{1cm} 265 \hspace{1cm} 270 \hspace{1cm}$

Leu Asn Glu Gly Ser Gln Ala Arg Pro Arg Ser Ser His Arg Tyr Phe 275 280 285

Leu Glu Arg Gly Leu Glu Ser Ala Thr Ser Leu 290 295

<210> 2

<211> 128

<212> PRT

<213> Mus sp.

<400> 2

Met Asp Leu Ser Glu Leu Glu Arg Asp Asn Thr Gly Arg Cys Arg Leu 1 5 10 15

Ser Ser Pro Val Pro Ala Val Cys Leu Leu Glu Pro Cys Val Leu Gly 20 25 30

Val Asp Glu Ala Gly Arg Gly Pro Val Leu Gly Pro Met Val Tyr Ala 35 40 45

Ile Cys Tyr Cys Pro Leu Ser Arg Leu Ala Asp Leu Glu Ala Leu Leu 50 55 60

Val Ala Asp Ser Leu Thr Leu Thr Glu Asn Glu Arg Glu Arg Leu Phe 65 70 75 80

Ala Leu Met Glu Glu Asp Gly Asp Phe Val Gly Trp Ala Leu Asp Val 85 90 95

Leu Ser Pro Asn Leu Ile Ser Thr Ser Met Leu Gly Arg Val Leu Tyr 100 105 110

Asn Leu Asn Ser Leu Ser His Asp Thr Ala Ala Gly Leu Ile Gln Tyr 115 120 125

<210> 3

<211> 307

<212> PRT

<213> Caenorhabditis elegans

<400> 3

Ser Leu Thr Val Leu Tyr Phe Ile Glu Arg Met Ser Leu Leu Cys Glu 1 5 10 15

Thr Glu Arg Ser Leu Thr Trp Asn Asn Phe Gly Asn Gly Ile Pro Cys 20 25 30

Val Leu Gly Ile Asp Glu Ala Gly Arg Gly Pro Val Leu Gly Pro Met 35 40 45

Val Tyr Ala Ala Ala Ile Ser Pro Leu Asp Gln Asn Val Glu Leu Leu 50 55 60

Asn Leu Gly Val Asp Asp Ser Leu Ala Leu Asn Glu Ala Leu Arg Glu 65 70 75 80

Glu Ile Phe Asn Leu Met Asn Glu Asp Glu Asp Ile Gln Gln Ile Ile 85 90 95

Ala Tyr Ala Leu Arg Cys Leu Ser Pro Glu Leu Ile Ser Cys Ser Met

100 105 110

Leu Leu Arg Gln Leu Tyr Ser Leu Asn Glu Val Ser His Glu Ala Ala 115 120 125

Ile Thr Leu Ile Arg Asp Ala Leu Ala Cys Asn Val Asn Val Val Glu 130 135 140

Glu Leu Leu Phe Pro Gly Ile Ser Ile Cys Val Thr Glu Leu Ala Asp 165 170 175

Ser Leu Phe Pro Ile Val Ser Ala Ala Ser Ile Ala Ala Leu Val Thr 180 185 190

Arg Asp Ser Arg Leu Arg Asn Trp Gln Phe Arg Glu Leu Asn Ile Leu 195 200 205

Val Pro Asp Ala Gly Tyr Gly Ser Gly Tyr Pro Gly Asp Pro Asn Thr 210 215 220

Leu Leu Phe Leu Gln Leu Ser Val Glu Pro Val Phe Gly Phe Cys Ser 225 230 235 240

Leu Val Arg Ser Ser Trp Leu Thr Ala Ser Thr Ile Val Glu Leu Arg 245 250 255

Cys Val Pro Gly Ser Trp Glu Asp Asp Glu Glu Glu Gly Leu Ser Gln 260 265 270

Ser Leu Arg Met Thr Ser Trp Met Val Pro Leu Asn Glu Thr Glu Val 275 280 285

Val Pro Leu Arg Asn Met Glu Ile Asn Leu Thr Leu Ile Val Ser Thr 290 295 300

Leu Phe Leu 305

<210> 4

<211> 307

<212> PRT

<213> Saccharomyces cerevisiae

<400> 4

Met Val Pro Pro Thr Val Glu Ala Ser Leu Glu Ser Pro Tyr Thr Leu 1 5 10 15

Ser Tyr Phe Ser Pro Val Pro Ser Ala Leu Leu Glu Gln Asn Asp Ser 20 25 30

Pro Ile Ile Met Gly Ile Asp Glu Ala Gly Arg Gly Pro Val Leu Gly 35 40 45

Pro Met Val Tyr Ala Val Ala Tyr Ser Thr Gln Leu Tyr Gln Asp Glu 50 55 60

Thr Ile Ile Pro Asn Tyr Glu Phe Asp Asp Ser Leu Leu Leu Thr Asp 65 70 75 80

Pro Ile Arg Arg Met Leu Phe Ser Leu Ile Tyr Gln Asp Asn Glu Glu 85 90 95

Leu Thr Gln Ile Gly Tyr Ala Thr Thr Cys Ile Thr Pro Leu Asp Ile 100 105 110

Ser Arg Gly Met Ser Leu Phe Pro Pro Thr Arg Asn Tyr Asn Leu Asn 115 120 125

Glu Gln Ala His Asp Val Thr Met Ala Leu Ile Asp Gly Val Ile Leu 130 135 140

Gln Asn Val Leu Leu Ser His Val Tyr Val Asp Thr Val Gly Pro Pro 145 150 155 160

Ala Ser Tyr Gln Leu Leu Glu Gln Arg Phe Pro Gly Val Leu Phe
165 170 175

Thr Val Ala Leu Leu Ala Asp Ser Leu Tyr Cys Met Val Ser Val Ala 180 185 190

Ser Val Val Ala Leu Val Thr Arg Asp Ile Leu Val Glu Ser Leu Leu 195 200 205

Arg Asp Pro Asp Glu Ile Leu Gly Ser Gly Tyr Pro Ser Asp Pro Leu 210 215 220

Thr Val Ala Trp Leu Leu Arg Asn Gln Thr Ser Leu Met Gly Trp Pro 225 230 235 240

Ala Asn Met Val Arg Phe Ser Trp Gln Thr Cys Gln Thr Leu Leu Asp 245 250 255

Asp Ala Ser Leu Asn Ser Ile Pro Ile Leu Trp Glu Glu Gln Tyr Met 260 265 270

Asp Ser Arg Leu Asn Ala Ala Gln Leu Thr Leu Gln Leu Gln Leu Gln 275 280 285

Met Val Ala Leu Pro Val Arg Arg Leu Arg Leu Arg Thr Leu Asp Asn 290 295 300

Trp Tyr Arg 305

<210> 5

<211> 198

<212> PRT

<213> Escherichia coli

<400> 5

Met Ile Glu Phe Val Tyr Pro His Thr Gln Leu Val Ala Gly Val Asp 1 5 10 15

Glu Val Gly Arg Gly Pro Leu Val Gly Ala Val Val Thr Ala Ala Val 20 25 30

Ile Leu Asp Pro Ala Arg Pro Ile Ala Gly Leu Asn Asp Ser Leu Leu 35 40 45

Leu Ser Glu Leu Arg Arg Leu Ala Leu Tyr Glu Glu Ile Leu Glu Leu 50 55 60

Ala Leu Ser Trp Ser Leu Gly Arg Ala Glu Pro His Glu Ile Asp Glu 65 70 75 80

Leu Asn Ile Leu His Ala Thr Met Leu Ala Met Gln Arg Ala Val Ala 85 90 95

Gly Leu His Ile Ala Pro Glu Tyr Val Leu Ile Asp Gly Asn Arg Cys
100 105 110

Pro Leu Leu Pro Met Pro Ala Met Ala Val Val Leu Gly Asp Ser Arg 115 120 125

Val Pro Glu Ile Ser Ala Ala Ser Ile Leu Ala Leu Val Thr Arg Asp 130 135 140 Ala Glu Met Ala Ala Leu Asp Ile Val Phe Pro Gln Tyr Gly Phe Ala 145 150 155 160

Gln His Leu Gly Tyr Pro Thr Ala Phe His Leu Glu Leu Leu Ala Glu 165 170 175

His Gly Ala Thr Glu His His Arg Arg Ser Phe Gly Pro Val Leu Arg 180 185 190

Ala Leu Gly Leu Ala Ser 195

<210> 6

<211> 286

<212> PRT

<213> Homo sapiens

<400> 6

Met Ser Trp Leu Leu Phe Leu Ala His Arg Val Ala Leu Ala Ala Leu 1 5 \cdot 10 15

Pro Cys Arg Arg Gly Ser Arg Gly Phe Gly Met Phe Tyr Ala Val Arg 20 25 30

Arg Gly Arg Leu Thr Gly Val Phe Leu Thr Trp Asn Glu Cys Arg Ala 35 40 45

Gln Val Asp Arg Phe Pro Ala Ala Arg Phe Leu Leu Phe Ala Thr Glu 50 55 60

Asp Glu Ala Trp Ala Phe Val Arg Leu Ser Ala Ser Pro Glu Val Ser 65 70 75 80

Glu Gly His Glu Asn Gln His Gly Gln Glu Ser Glu Ala Leu Pro Gly 85 90 95

Leu Arg Leu Arg Glu Pro Leu Asp Gly Asp Gly His Glu Ser Ala Gln
100 105 110

Pro Tyr Ala Leu His Met Leu Pro Ser Val Glu Pro Ala Pro Pro Val 115 120 125

Ser Arg Asp Thr Phe Ser Tyr Met Gly Asp Phe Val Val Tyr Thr 130 135 140

Gly Val Tyr Trp Gly Pro Gly His Pro Leu Asn Val Gly Ile Arg Leu 165 170 175

Pro Gly Arg Gln Thr Asn Gln Arg Ala Glu Ile His Ala Ala Cys Leu 180 185 190

Ala Ile Glu Gln Ala Leu Thr Gln Asn Ile Asn Leu Leu Val Leu Tyr 195 200 205

Thr Asp Ser Met Phe Thr Ile Asn Gly Ile Thr Asn Trp Val Gln Gly 210 215 220

Trp Leu Leu Asn Gly Trp Leu Thr Ser Ala Gly Leu Glu Val Ile Asn 225 230 235 240

Leu Glu Asp Phe Val Ala Leu Glu Arg Leu Thr Gln Gly Met Asp Ile 245 250 255

Gln Trp Met His Val Pro Gly His Ser Gly Phe Ile Gly Asn Glu Glu 260 265 270

Ala Asp Arg Leu Ala Arg Glu Gly Ala Leu Gln Ser Glu Asp 275 280 285

<210> 7

<211> 286

<212> PRT

<213> Homo sapiens

<400> 7

Met Ser Trp Leu Leu Phe Leu Ala His Arg Val Ala Leu Ala Ala Leu 1 5 10 15

Pro Cys Arg Arg Gly Ser Arg Gly Phe Gly Met Phe Tyr Ala Val Arg 20 25 30

Arg Gly Arg Leu Thr Gly Val Phe Leu Thr Trp Asn Glu Cys Arg Ala 35 40 45

Gln Val Asp Arg Phe Pro Ala Ala Arg Phe Leu Leu Phe Ala Thr Glu 50 60

Asp Glu Ala Trp Ala Phe Val Arg Leu Ser Ala Ser Pro Glu Val Ser 65 70 75 80

Glu Gly His Glu Asn Gln His Gly Gln Glu Ser Glu Ala Leu Ala Ser

Leu Arg Leu Arg Glu Pro Leu Asp Gly Asp Gly His Glu Ser Ala Glu 100 105

Pro Tyr Ala Leu His Met Leu Pro Ser Val Glu Pro Ala Pro Pro Val 115 120

Ser Arg Asp Thr Phe Ser Tyr Met Gly Asp Phe Val Val Val Tyr Thr 130 135

Asp Gly Cys Cys Ser Ser Asn Gly Arg Arg Pro Arg Ala Gly Ile 145 150 155

Gly Val Tyr Trp Gly Pro Gly His Pro Leu Asn Val Gly Ile Arg Leu 165 170

Pro Gly Arg Gln Thr Asn Gln Arg Ala Glu Ile His Ala Ala Cys Leu 180 185 190

Ala Ile Glu Gln Ala Leu Thr Gln Asn Ile Asn Leu Leu Val Leu Tyr

Thr Asp Ser Met Phe Thr Ile Asn Gly Ile Thr Asn Trp Val Gln Gly 215

Trp Leu Leu Asn Gly Trp Leu Thr Ser Ala Gly Leu Glu Val Ile Asn 225 230 235

Leu Glu Asp Phe Val Ala Leu Glu Arg Leu Thr Gln Gly Met Asp Ile 245

Gln Trp Met His Val Pro Gly His Ser Gly Phe Ile Gly Asn Glu Glu

Ala Asp Arg Leu Ala Arg Glu Gly Ala Leu Gln Ser Glu Asp 280

<210> 8

<211> 286 <212> PRT <213> Homo sapiens

<400> 8

Met Ser Trp Phe Leu Phe Leu Ala His Arg Val Ala Leu Ala Ala Leu 1 5 10 15

Pro Cys Arg Arg Gly Ser Arg Gly Phe Gly Met Phe Tyr Ala Val Arg 20 25 30

Arg Gly Arg Leu Thr Gly Val Phe Leu Thr Trp Asn Glu Cys Arg Ala 35 40 45

Gln Val Asp Arg Phe Pro Ala Ala Arg Phe Leu Leu Phe Ala Thr Glu 50 55 60

Asp Glu Ala Trp Ala Phe Val Arg Leu Ser Ala Ser Pro Glu Val Ser 65 70 75 80

Glu Gly His Glu Asn Gln His Gly Gln Glu Ser Glu Ala Leu Ala Ser 85 90 95

Leu Arg Leu Arg Glu Pro Leu Asp Gly Asp Gly His Glu Ser Ala Glu 100 105 110

Pro Tyr Ala Leu His Met Leu Pro Ser Val Glu Pro Ala Pro Pro Val 115 120 125

Ser Arg Asp Thr Phe Ser Tyr Met Gly Asp Phe Val Val Tyr Thr 130 135 140

Gly Val Tyr Trp Gly Pro Gly His Pro Leu Asn Val Gly Ile Arg Leu 165 170 175

Pro Gly Arg Gln Thr Asn Gln Arg Ala Glu Ile His Ala Ala Cys Leu 180 185 190

Ala Ile Glu Gln Ala Leu Thr Gln Asn Ile Asn Leu Leu Val Leu Tyr 195 200 205

Thr Asp Ser Met Phe Thr Ile Asn Gly Ile Thr Asn Trp Val Gln Gly 210 215 220

Trp Leu Leu Asn Gly Trp Leu Thr Ser Ala Gly Leu Glu Val Ile Asn 225 230 235 240

Leu Glu Asp Phe Val Ala Leu Glu Arg Leu Thr Gln Gly Met Asp Ile

245 250 255

Gln Trp Met His Val Pro Gly His Ser Gly Phe Ile Gly Asn Glu Glu 260 265 270

Ala Asp Arg Leu Ala Arg Glu Gly Ala Leu Gln Ser Glu Asp 275 280 285

<210> 9

<211> 286

<212> PRT

<213> Homo sapiens

<400> 9

Met Ser Trp Leu Leu Phe Leu Ala His Arg Val Ala Leu Ala Ala Leu 1 5 10 15

Pro Cys Arg Arg Gly Ser Arg Gly Phe Gly Met Phe Tyr Ala Val Arg 20 25 30

Arg Gly Arg Leu Thr Gly Val Phe Leu Thr Trp Asn Glu Cys Arg Ala 35 40 45

Gln Val Asp Arg Phe Pro Ala Ala Arg Phe Leu Leu Phe Ala Thr Glu 50 55 60

Asp Glu Ala Trp Ala Phe Val Arg Leu Ser Ala Ser Pro Glu Val Ser 65 70 75 80

Glu Gly His Glu Asn Gln His Gly Arg Glu Ser Glu Ala Leu Ala Ser 85 90 95

Leu Arg Leu Arg Glu Pro Leu Asp Gly Asp Gly His Glu Ser Ala Glu 100 105 110

Pro Tyr Ala Leu His Met Leu Pro Ser Val Glu Pro Ala Pro Pro Val 115 120 125

Ser Arg Asp Thr Phe Ser Tyr Met Gly Asp Phe Val Val Tyr Thr 130 135 140

Asp Gly Cys Cys Ser Ser Asn Gly Arg Arg Pro Arg Ala Gly Ile 145 150 155 160

Gly Val Tyr Trp Gly Pro Gly His Pro Leu Asn Val Gly Ile Arg Leu 165 170 175

Pro Gly Arg Gln Thr Asn Gln Arg Ala Glu Ile His Ala Ala Cys Leu 180 185 190

Ala Ile Glu Gln Ala Leu Thr Gln Asn Ile Asn Leu Leu Val Leu Tyr 195 200 205

Thr Asp Ser Met Phe Thr Ile Asn Gly Ile Thr Asn Trp Val Arg Gly 210 215 220

Trp Leu Leu Asn Gly Trp Leu Thr Ser Ala Gly Leu Glu Val Ile Asn 225 230 235 240

Leu Glu Asp Phe Val Ala Leu Glu Arg Leu Thr Gln Gly Met Asp Ile 245 250 255

Gln Trp Met His Val Pro Gly His Ser Gly Phe Ile Gly Asn Glu Glu 260 265 270

Ala Asp Arg Leu Ala Arg Glu Gly Ala Leu Gln Ser Glu Asp 275 280 285

<210> 10

<211> 299

<212> PRT

<213> Homo sapiens

<400> 10

Met Asp Leu Ser Glu Leu Glu Arg Asp Asn Thr Gly Arg Cys Arg Leu 1 5 10 15

Ser Ser Pro Val Pro Ala Val Cys Arg Leu Glu Pro Cys Val Leu Gly
20 25 30

Val Asp Glu Ala Gly Arg Gly Pro Val Leu Gly Pro Met Val Tyr Ala 35 40 45

Ile Cys Tyr Cys Pro Leu Pro Arg Leu Ala Asp Leu Glu Ala Leu Leu 50 55 60

Val Ala Asp Ser Leu Thr Leu Leu Glu Ser Glu Arg Glu Arg Leu Phe 70 75 80

Ala Leu Met Glu Asp Thr Asp Phe Val Gly Trp Ala Leu Asp Val Leu 85 90 95

Ser Pro Asn Leu Ile Ser Thr Ser Met Leu Gly Arg Val Leu Tyr Asn

100 105 110

Leu Asn Ser Leu Ser His Asp Thr Ala Thr Gly Leu Ile Gln Tyr Ala 115 120 125

Leu Asp Gln Gly Val Asn Val Thr Gln Val Phe Val Asp Thr Val Gly 130 135 140

Glu Val Thr Val Leu Ala Leu Ala Asp Ala Leu Tyr Pro Val Val Ser 165 170 175

Ala Ala Ser Ile Cys Ala Leu Val Ala Arg Asp Gln Ala Val Leu Leu 180 185 190

Trp Gln Phe Val Glu Leu Leu Gln Asp Leu Asp Thr Asp Tyr Gly Ser 195 200 205

Gly Tyr Pro Asn Asp Pro Leu Thr Leu Ala Trp Leu Leu Glu His Val 210 215 220

Glu Pro Val Phe Gly Phe Pro Gln Phe Val Arg Phe Ser Trp Arg Thr 225 230 235 240

Ala Gln Thr Ile Leu Glu Leu Glu Ala Glu Asp Val Ile Trp Glu Asp 245 250 255

Ser Ala Ser Glu Asn Gln Glu Gly Leu Arg Leu Ile Thr Ser Tyr Phe 260 265 270

Leu Asn Glu Gly Ser Gln Ala Arg Pro Arg Ser Ser His Arg Tyr Phe 275 280 285

Leu Glu Arg Gly Leu Glu Ser Ala Thr Ser Leu 290 295

<210> 11

<211> 285

<212> PRT

<213> Mus sp.

<400> 11

Met Arg Trp Leu Leu Pro Leu Ser Arg Thr Val Thr Leu Ala Val Val 1 5 10 15

- Arg Leu Arg Arg Gly Ile Cys Gly Leu Gly Met Phe Tyr Ala Val Arg 20 25 30
- Arg Gly Arg Arg Thr Gly Val Phe Leu Ser Trp Ser Glu Cys Leu Ala 35 40 45
- Gln Val Asp Arg Phe Pro Ala Ala Arg Phe Leu Leu Phe Ala Thr Glu 50 55 60
- Asp Glu Ala Trp Ala Phe Val Arg Ser Ser Ser Ser Pro Asp Gly Ser 65 70 75 80
- Leu Gly Gln Glu Ser Ala His Glu Gln Leu Ser Gln Ala Leu Thr Ser 85 90 95
- Leu Arg Pro Arg Glu Pro Leu Gly Glu Glu Glu Leu Pro Glu Pro 100 105 110
- Gly Pro Leu His Thr Arg Gln Asp Thr Glu Pro Ala Ala Val Val Ser 115 120 125
- Leu Asp Thr Phe Ser Tyr Met Gly Glu Ser Val Ile Val Tyr Thr Asp 130 135 140
- Val Tyr Trp Gly Pro Gly His Pro Leu Asn Val Gly Ile Arg Leu Pro 165 170 175
- Gly Arg Gln Thr Asn Gln Arg Ala Glu Ile His Ala Ala Cys Leu Ala 180 185 190
- Ile Met Gln Ala Leu Ala Gln Asn Ile Ser Leu Leu Val Leu Tyr Thr 195 200 205
- Asp Ser Met Phe Thr Ile Asn Gly Ile Thr Asn Trp Val Gln Gly Trp 210 215 220
- Leu Leu Asn Gly Trp Arg Thr Ser Thr Gly Leu Asp Val Ile Asn Leu 225 230 235 240
- Glu Asp Phe Met Glu Leu Asp Glu Leu Thr Gln Gly Met Asp Ile Gln
 245 250 255

Trp Met His Ile Pro Gly His Ser Gly Phe Val Gly Asn Glu Glu Ala 265 260

Asp Arg Leu Ala Arg Glu Gly Ala Leu Gln Ser Glu Asp 280

<210> 12 <211> 1131 <212> DNA

<213> Homo sapiens

<400> 12

cgcgcctgca	gtattagttc	ttgcagctgg	tggtggcggc	tgaggcggca	tggatctcag	60
cgagctggag	agagacaata	caggccgctg	tcgcctgagt	tcgcctgtgc	ccgcggtgtg	120
ccgcaaggag	ccttgcgtcc	tgggcgtcga	tgaggcgggc	aggggccccg	tgctgggccc	180
catggtctac	gccatctgtt	attgtcccct	gcctcgcctg	gcagatctgg	aggcgctgaa	240
agtggcagac	tcaaagaccc	tattggagag	cgagcgggaa	aggctgtttg	cgaaaatgga	300
ggacacggac	tttgtcggct	gggcgctgga	tgtgctgtct	ccaaacctca	tctctaccag	360
catgcttggg	tgggtcaaat	acaacctgaa	ctccctgtca	catgatacag	ccactgggct	420
tatacagtat	gcattggacc	agggcgtgaa	cgtcacccag	gtattcgtgg	acaccgtagg	480
gatgccagag	acataccagg	cgcggctgca	gcaaagtttt	cccgggattg	aggtgacggt	540
caaggccaaa	gcagatgccc	tctacccggt	ggttagtgct	gccagcatct	gtgccaaggt	600
ggcccgggac	caggccgtga	agaaatggca	attcgtggag	aaactgcagg	acttggatac	660
tgattatggc	tcaggctacc	ccaatgatcc	caagacaaaa	gcgtggttga	aggagcacgt	720
ggagcctgtg	ttcggcttcc	cccagtttgt	ccggttcagc	tggcgcacgg	cccagaccat	780
cctggagaaa	gaggcggaag	atgttatatg	ggaggactca	gcatccgaga	atcaggaggg	840
actcaggaag	atcacatcct	acttcctcaa	tgaagggtcc	caagcccgtc	cccgttcttc	900
ccaccgatat	ttcctggaac	gcggcctgga	gtcagcaacc	agcctctagc	agctgcctct	960
acgcgctcta	cctgcttccc	caacccagac	attaaaattg	tttaaggaga	accacacgta	1020
ggggatgtac	ttttgggaca	gaagcaaggt	gggagtgtgc	tctgcagccg	ggtccagcta	1080
cttccttttg	gaaccttaaa	tagaatgggt	gttggttgat	aaaaaaaaa	a	1131

<210> 13

<211> 20 <212> DNA <213> Artificial sequence

<220>

<223> Synthetic

<400> agcaggo	13 egec gettegagge	20
<210>	14	
<211>	26	
<212>		
	Artificial sequence	
<220>	Synthetic	
(223)	Synchecte	
<400>	14 .	
cccgct	cctg cagtattagt tettge	26
<210>	15	
<211>	25	
<212>		
<213>	Artificial sequence	
<220>		
<223>	Synthetic	
<400>	15	
	ctgg tggtggcggc tgagg	25
<210>	16	
<211>		
<212>		
	Artificial sequence	
4220s		
<220> <223>	Synthetic	
	-7	
<400>	16	
tccaata	aggg tetttgagte tgecae	26
<210>	17	
<211>	25	
	DNA Artificial sequence	
(213)	Altilitial sequence	
<220>		
<223>	Synthetic	
<400>	17	
	cage geetecagat etgee	25
<210>	18	
<211>	26	
<212>		
	Artificial sequence	
<220> <223>	Synthetic	
~~ <i>~</i>	oynence10	
-400>	10	

gcgaggcagg ggacaataac agatgg 26			
<210><211><211><212><213>	17		
<220>	Synthetic		
<400> gggcgc	19 ecgtc ggtgtgg	17	
<210><211><211><212><213>	20		
<220> <223>	Synthetic		
<400> cgcctc	20 agcc gccaccacca	20	
<210><211><212><213>	20		
<220> <223>	Synthetic		
<400> cacagg	21	20	
<210><211><211><212><213>	20		
<220> <223>	Synthetic		
<400> ggacaa	22 taac agatggcgta	20	
<210><211><211><212><213>			
<220> <223>	Synthetic		
<400> cccgct	23 cgct ctccaatagg	20	

<210>	24	
<211>		
<212>		
<213>	Artificial sequence	
<220>		
<223>	Synthetic	
<400>	24	
cccagc	cgac aaagtccgtg	20
<210>	25	
<211>	20	
<212>		
<213>	Artificial sequence	
<220>		
<223>	Synthetic	
<400>	25	
cggtgt	ccac gaatacctgg	20
<210>		
<211>	20	
<212>	DNA	
<213>	Artificial sequence	
<220>		
	Synthetic	
	Synthetic	
<223> <400>		20
<223> <400>	26	20
<223> <400>	26	20
<223> <400> cgcgcc	26 tggt atgtetetgg 27	20
<223> <400> cgcgcc	26 tggt atgtetetgg 27	20
<223> <400> cgcgcc <210> <211> <212>	26 tggt atgtetetgg 27 20 DNA	20
<223> <400> cgcgcc	26 tggt atgtetetgg 27 20 DNA	20
<223> <400> cgcgcc <210> <211> <212> <213>	26 tggt atgtetetgg 27 20 DNA	20
<223> <400> cgcgcc <210> <211> <212> <213> <220>	26 tggt atgtctctgg 27 20 DNA Artificial sequence	20
<223> <400> cgcgcc <210> <211> <212> <213> <220>	26 tggt atgtetetgg 27 20 DNA	20
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic	20
<223> <400> cgcgcc <210> <211> <212> <213> <220>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic	20
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic	20
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtagae	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg	
<223> <400> cgcgcc <210> <211> <212> <213> <400> <ggtagac <210=""></ggtagac>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtagac <210> <211>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg 28 20	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtaga <210> <211> <212>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg 28 20 DNA	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtagac <210> <211>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg 28 20	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtagac <210> <211> <212> <213>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg 28 20 DNA	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtaga <210> <211> <212> <213> <	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg 28 20 DNA Artificial sequence	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtagac <210> <211> <212> <213>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg 28 20 DNA	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtaga <210> <211> <212> <213> <	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 tggc atctgctttg 28 20 DNA Artificial sequence Synthetic	
<223> <400> cgcgcc <210> <211> <212> <213> <220> <223> <400> ggtaga <210> <211> <212> <213> <400> <311> <212> <400> <400>	26 tggt atgtctctgg 27 20 DNA Artificial sequence Synthetic 27 gggc atctgctttg 28 20 DNA Artificial sequence	

<210>	29	
<211>	20	
<212>	DNA	
<213>	Artificial sequence	
	-	
<220>	·	
<223>	Synthetic	
12237	Synthetic	
.400-	20	
<400>	29	
cagttt	ctcc acgaattgcc	20
<210>	30	
<211>	20	
<212>	DNA	
<213>	Artificial sequence	
	-	
<220>		
<223>	Synthetic	
\2237	Synchectic .	
<400>	30	
LLLLGL	cttg ggatcattgg	20
	•	
<210>		
<211>	20	
<212>	DNA	
<213>	Artificial sequence	
	· · · · · · · · · · · · · · · · · · ·	
<220>		
<223>	Synthetic	
12232	o, neneere	
<400>	31	
		~ ~
agetga	accg gacaaactgg	20
<210>	32	
<211>		
<212>	DNA	
<213>	Artificial sequence	
	-	
<220>		
<223>	Synthetic	
12207	5,110110020	
<400>	32	
		2.0
CCLCLL	toto caggatggto	20
<210>	33	
<211>	20	
<212>	DNA	
<213>	Artificial sequence	
	-	
<220>		
<223>	Synthetic	
	-4	
<400>	33	
		2.0
acticca	ggcc gcgttccagg	20

```
<210> 34
<211> 20
<212> DNA
<213> Artificial sequence
<220>
<223> Synthetic
<400> 34
cctacgtgtg gttctcctta
                                                                 20
<210> 35
<211> 20
<220>
<223> Synthetic
<400> 35
                                                                 20
gcacactccc accttgcttc
<210> 36
<211> 20
<212> DNA
<213> Artificial sequence
<220>
<223> Synthetic
<400> 36
                                                                 20
caaaaggaag tagctggacc
<210> 37
<211> 20
<212> DNA
<213> Artificial sequence
<220>
<223> Synthetic
<400> 37
aaacaatttt aatgtctggg
                                                                 20
<210> 38
<211> 20
<212> DNA
<213> Artificial sequence
<220>
<223> Synthetic
<400> 38
aattttaatg tctgggttgg
                                                                 20
```

<210> 39

<211> 20 <212> DNA <213> Artificial sequence

<220>

<223> Synthetic

<400> 39

ccttaaacaa ttttaatgtc

20